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1 M E S L L L P R L E C N G A I 16

63 TCA GCT CAC CGC AAC CTC CGC CTC CCG GGT TCA AGC GAT TCT CCT GCC TCA GCC TCC CCA 122
17 S A H R N L R L P G S S D S P A S A S P 36

123 GTA GCT GGG ATT ACA GGC ATG TGC ACC CAC GCT CGG CTA ATT TTG TAT TTT TTT TTA GTA 182
37 V A G I T G M C T H A R L I L Y F F L V 56

183 GAG ATG GAG TTT CTC CAT GTT GGT CAG GCT GGT CTC GAA CTC CCG ACC TCA GAT GAT CCC 242
57 E M E F L H V G Q A G L E L P T S D D P 76

243 TCC GTC TCG GCC TCC CAA AGT GCT AGA TAC AGG ACT GGC CAC CAT GCC CGG CTC TGC CTG 302
77 S V S A S Q S A R Y R T G H H A R L C L 96

303 GCT AAT TTT TGT GGT AGA AAC AGG GTT TCA CTG ATG TGC CCA AGC TGG TCT CCT GAG CTC 362
97 A N F C G R N R V S L M C P S W S P E L 116

363 AAG CAG TCC ACC TGC CTC AGC CTC CCA AAG TGC TGG GAT TAC AGG CGT GCA GCC GTG CCT 422
117 K Q S T C L S L P K C W D Y R R A A V P 136

423 GGC CTT TTT ATT TTA TTT TTT TTA AGA CAC AGG TGT GGG ACT GGT ACC GAG GAT GAT GAG 482
137 G L F I L F F L R H R C P T L T O D E V 156

483 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 542
157 Q W C D H S S L Q P S T P E I K H P P A 176

543 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 602
177 S A S Q V A G T K D M H H Y T W L I F I 196

603 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 662
197 F I F N F L R Q S L N S V T Q A G V Q W 216

663 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 722
217 R N L G S L Q P L P P G F K L F S C P S 236

723 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 782
237 L L S S W D Y R R P P R L A N F F V F L 256

783 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 842
257 V E M G F T M F A R L I L I S G P C D L 276

843 GAG TCC TGT GAT GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG GAG 902
277 P A S A S Q S A G I T G V S H H A R L I 296

903 TTT AAT TTT TGT TTG TTT GAA ATG GAA TCT CAC TCT GTT ACC CAG GCT GGA GTG CAA TGG 962
297 F N F C L F E M E S H S V T Q A G V Q W 316

963 CCA AAT CTC GGC TCA CTG CAA CCT CTG CCT CCC GGG CTC AAG CGA TTC TCC TGT CTC AGC 1022
317 P N L G S L Q P L P P G L K R F S C L S 336

1023 CTC CCA AGC AGC TGG GAT TAC GGG CAC CTG CCA CCA CAC CCC GCT AAT TTT TGT ATT TTC 1082
337 L P S S W D Y G H L P P H P A N F C I F 356

1083 ATT AGA GGC GGG GTT TCA CCA TAT TTG TCA GGC TGG TCT CAA ACT CCT GAC CTC AGG tgac 1143
357 I R G G V S P Y L S G W S Q T P D L R 375

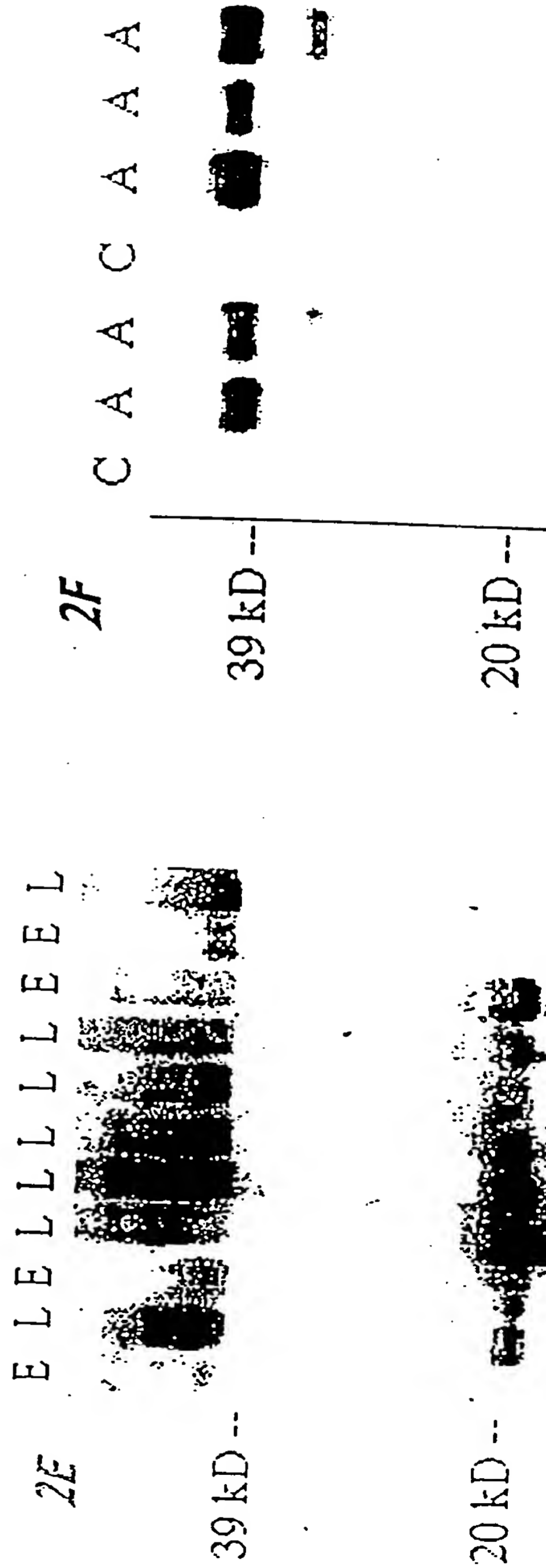
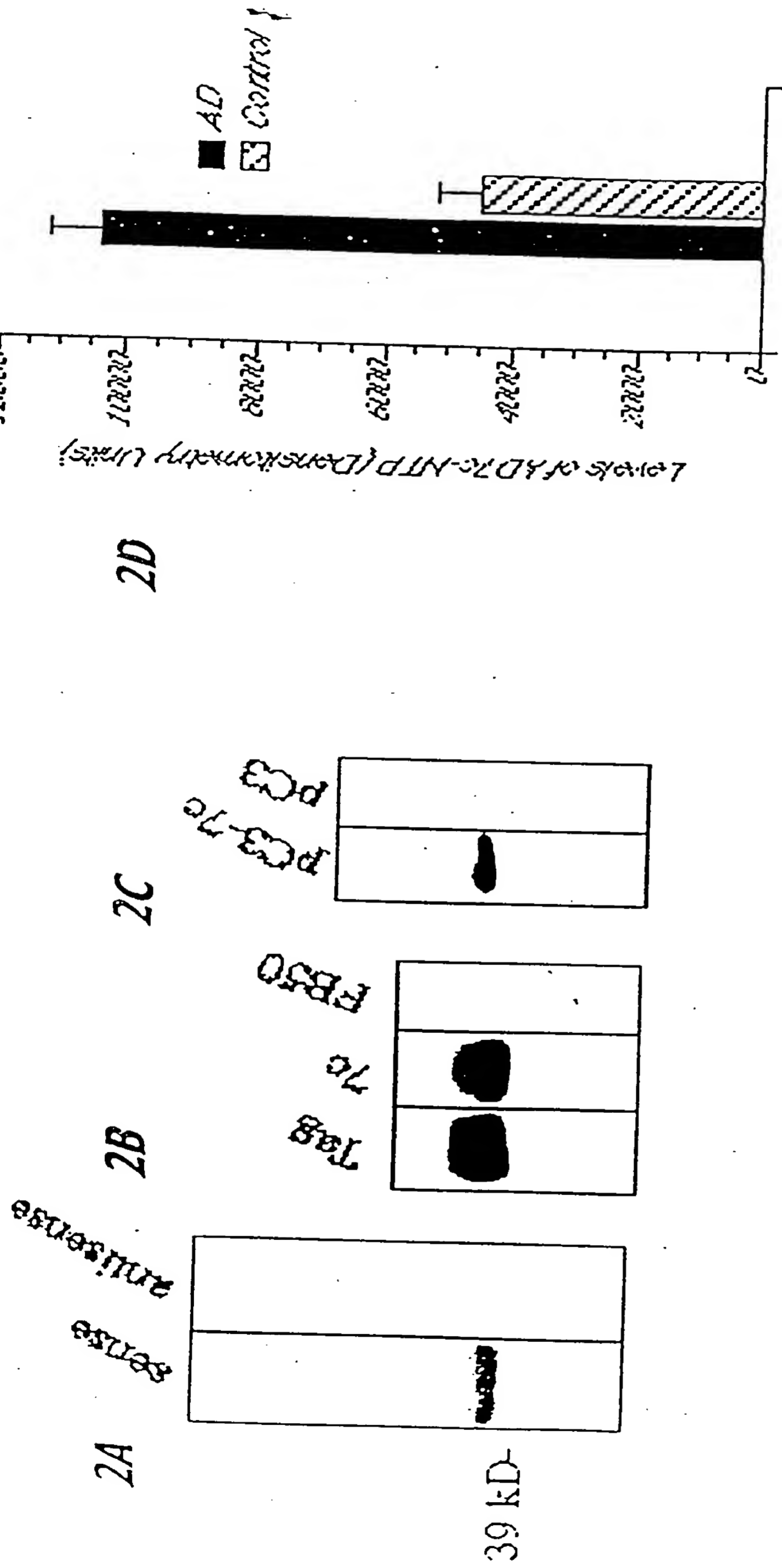
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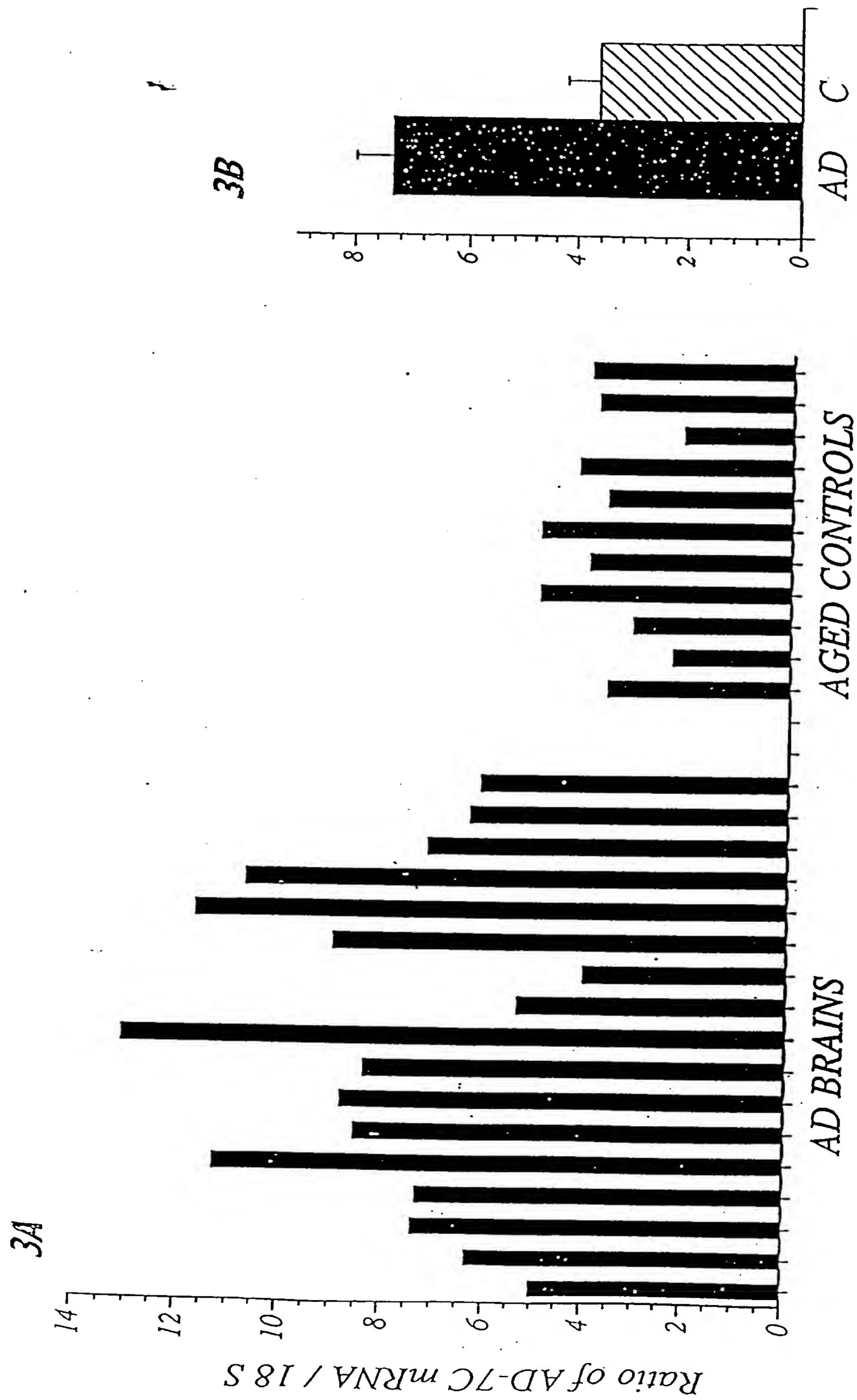
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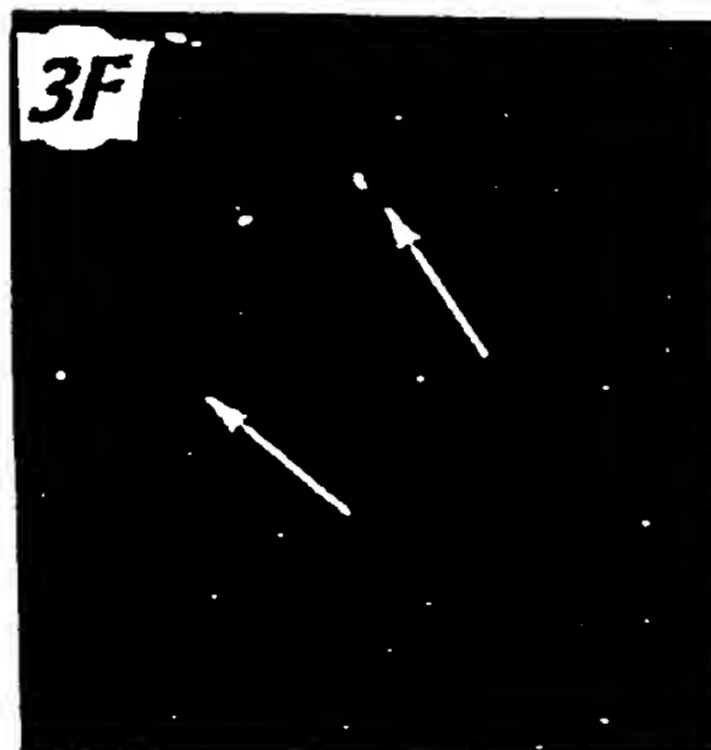
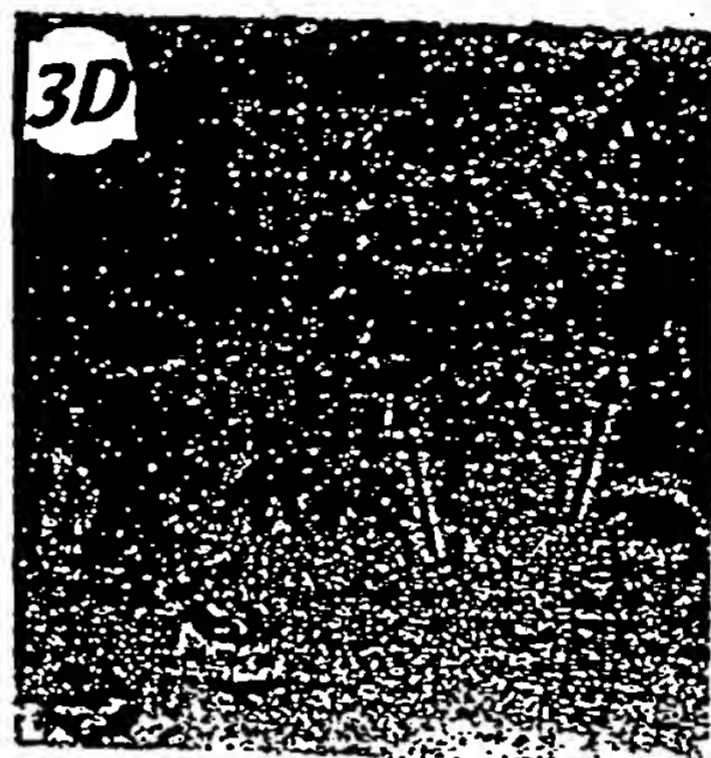
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FIG. 1



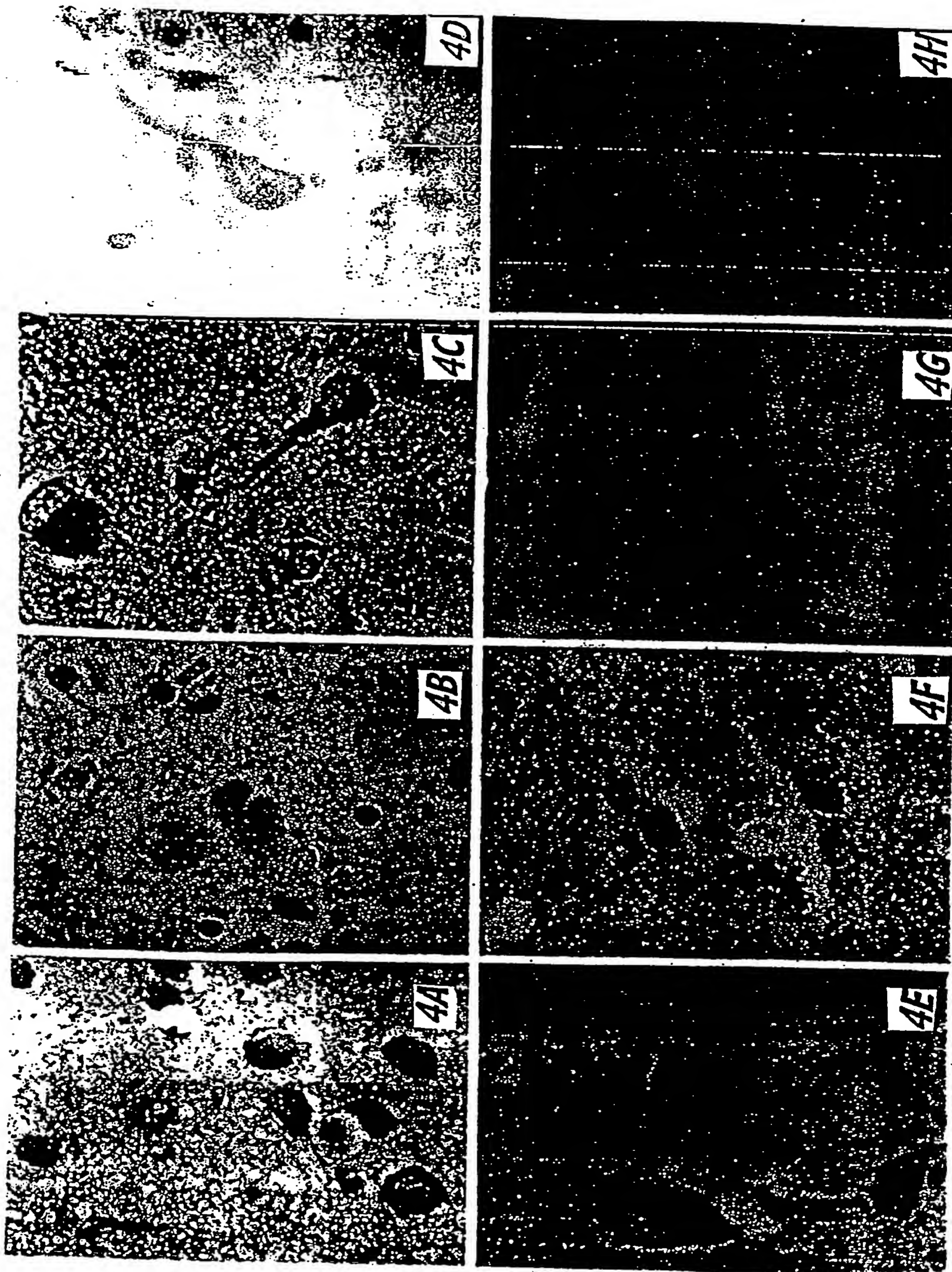


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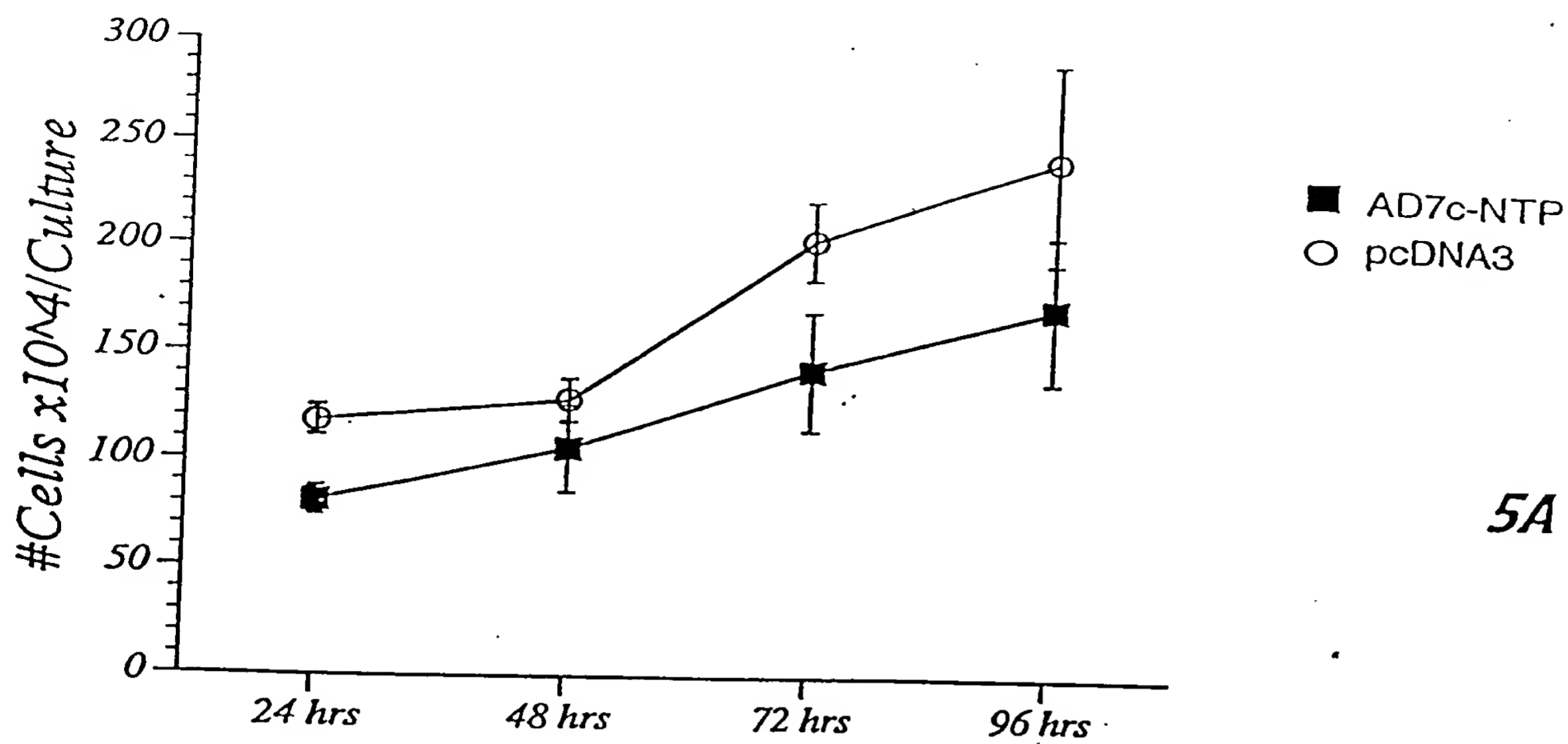


FIGS. 3C-3F

FIGS. 4A-4H

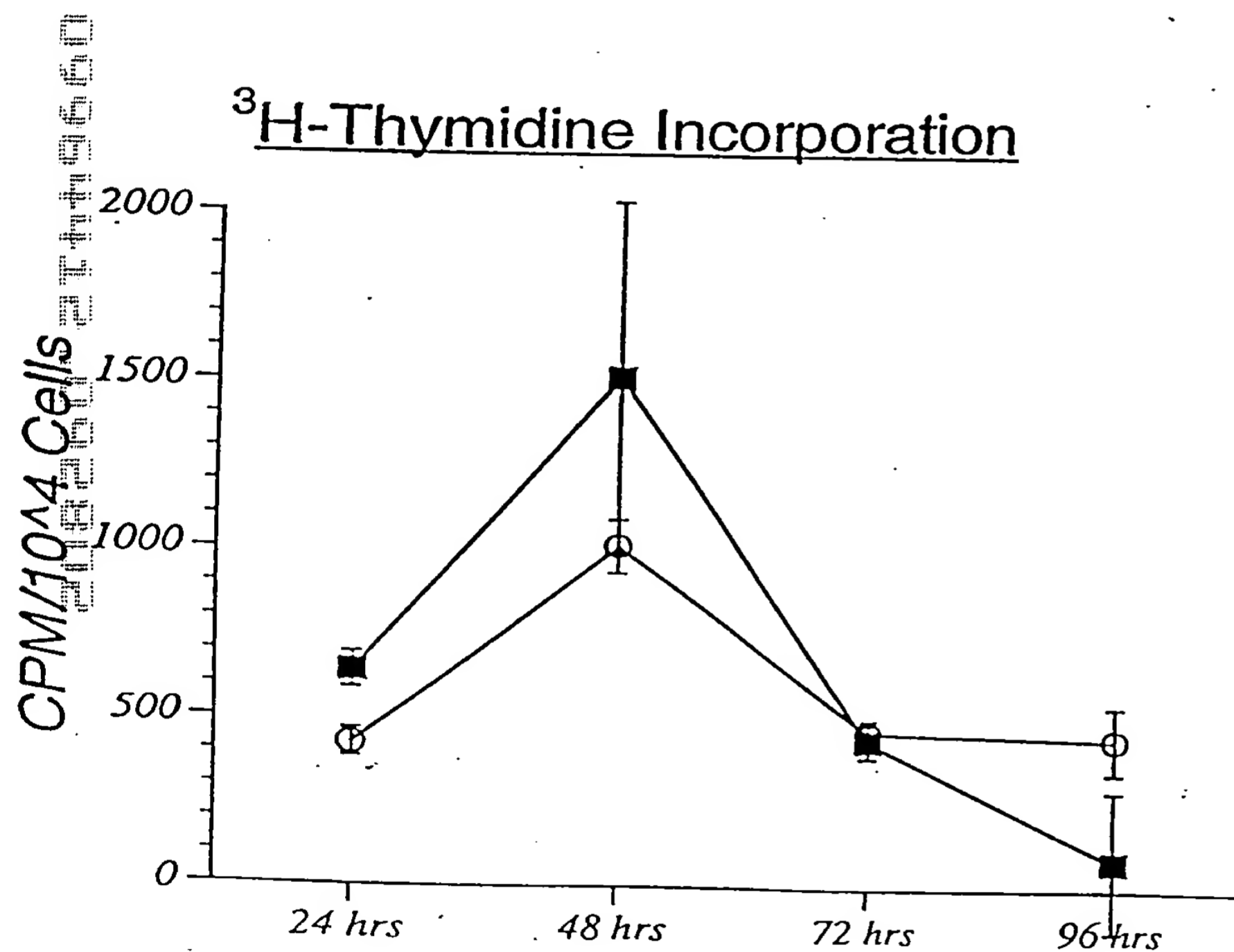


Growth of SH-Sy5y Cells



5A

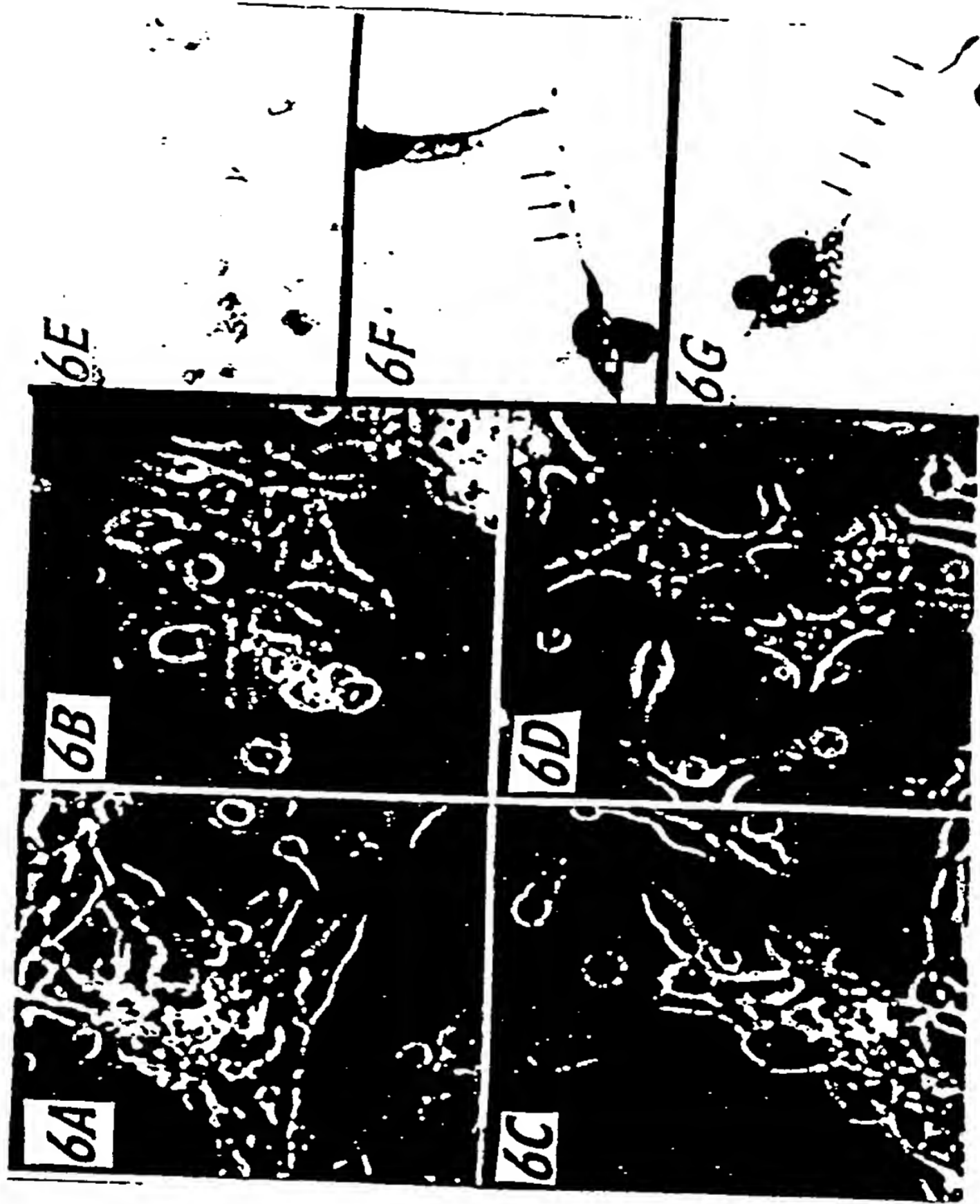
3 H-Thymidine Incorporation

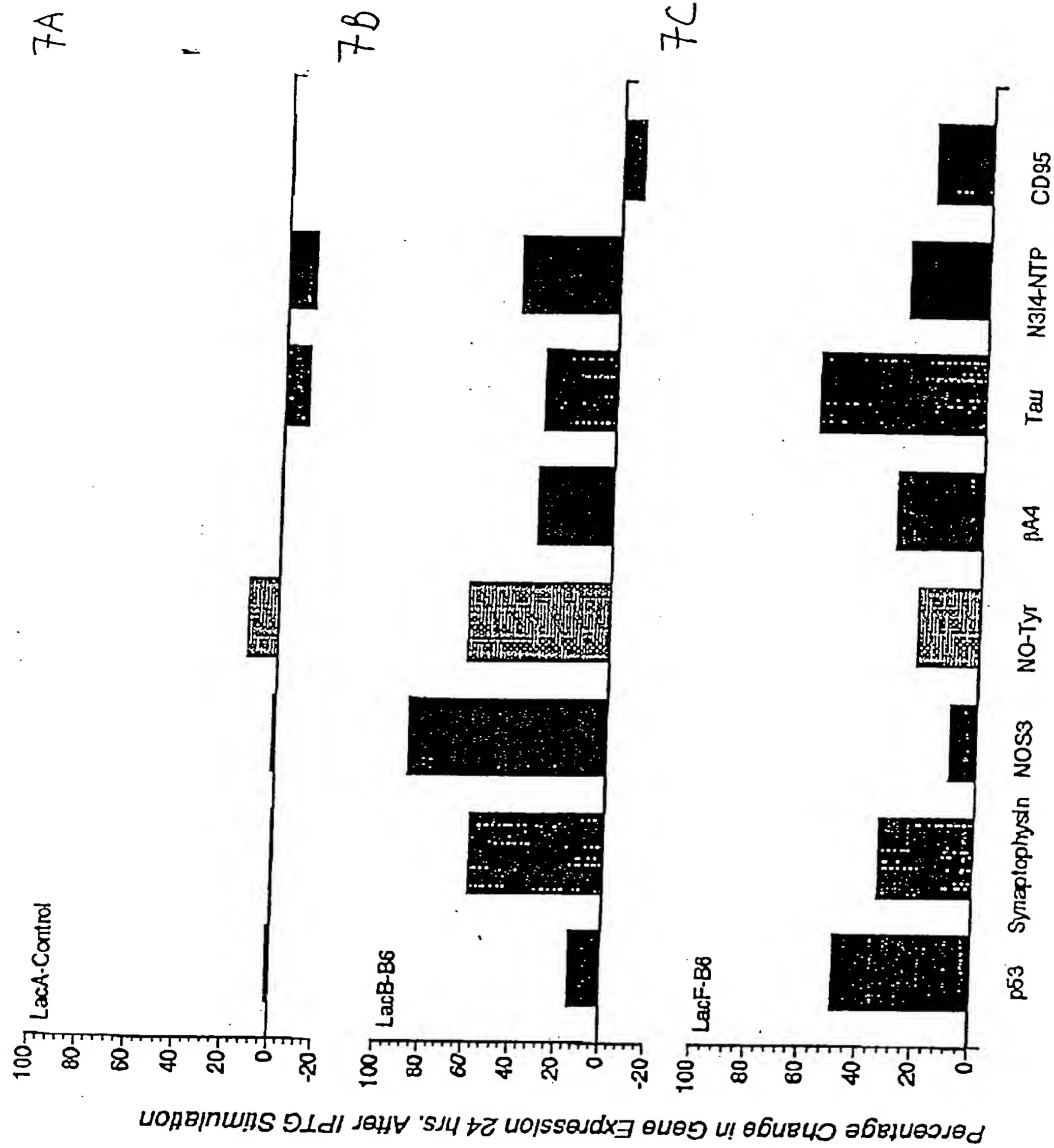


5B

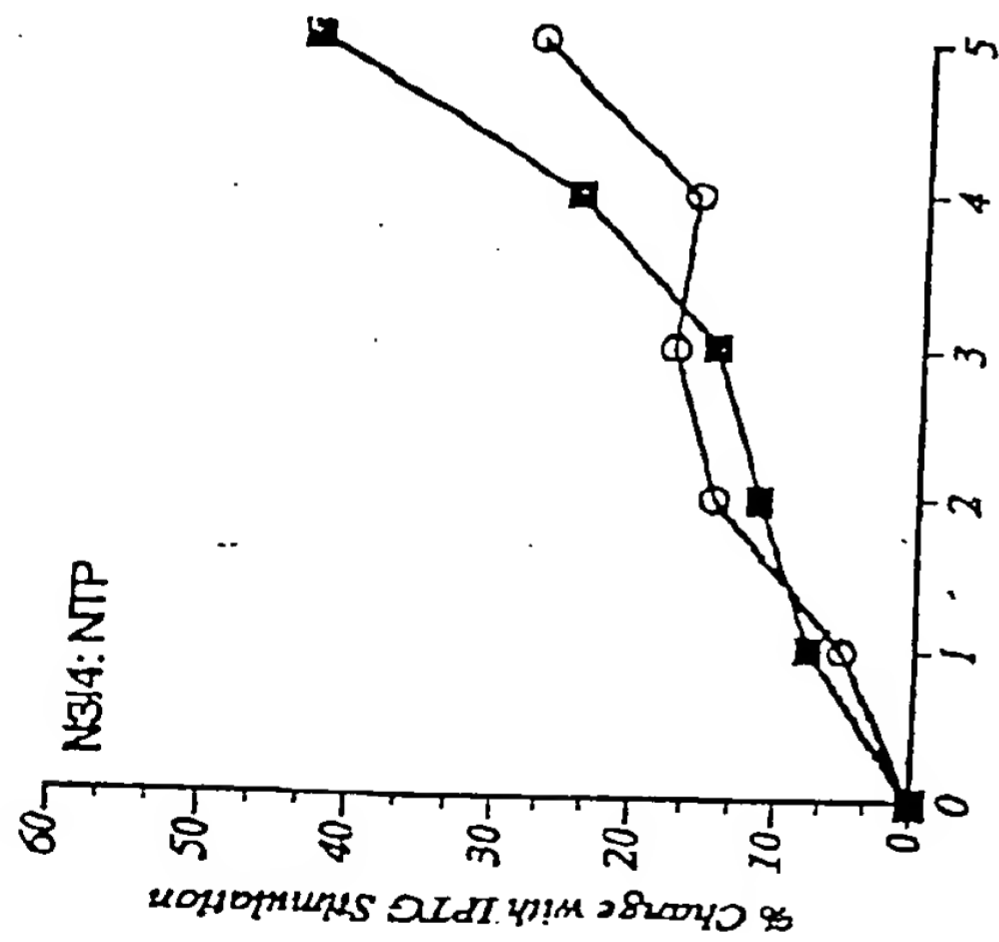
FIGS. 5A-5B

FIGS. 6A-6G

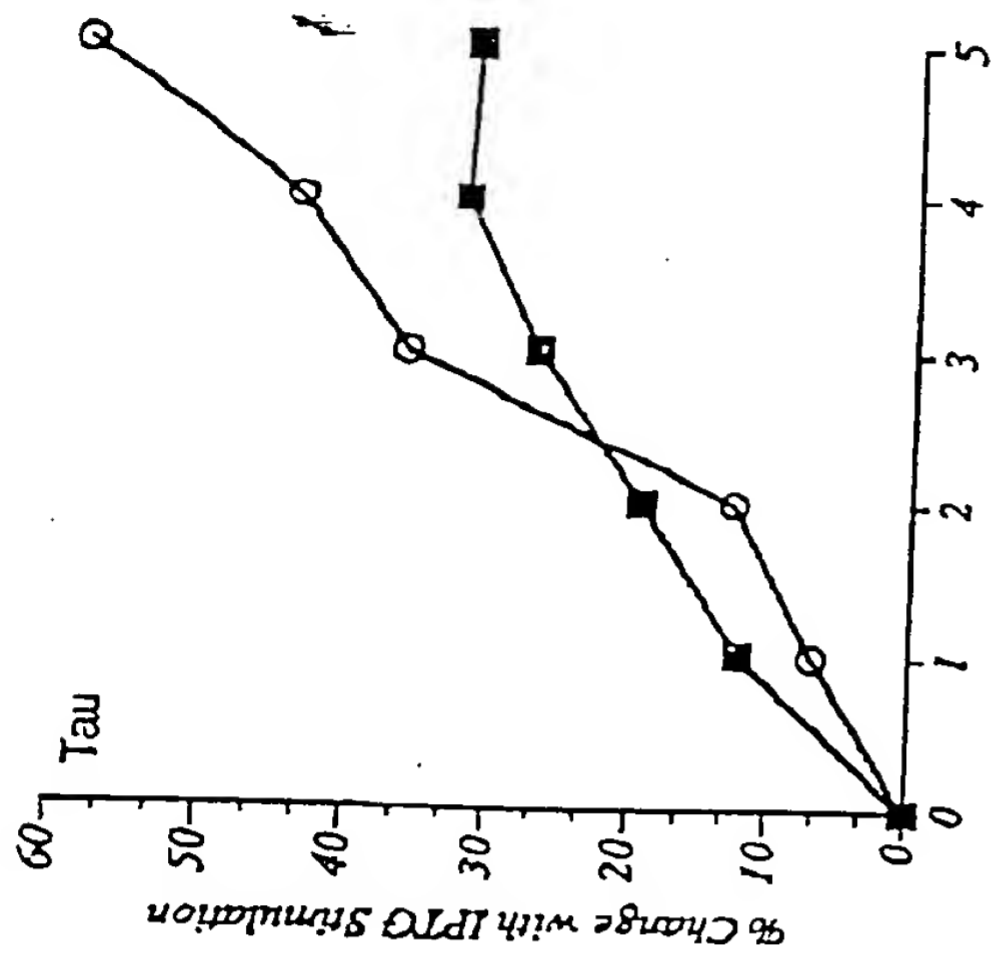




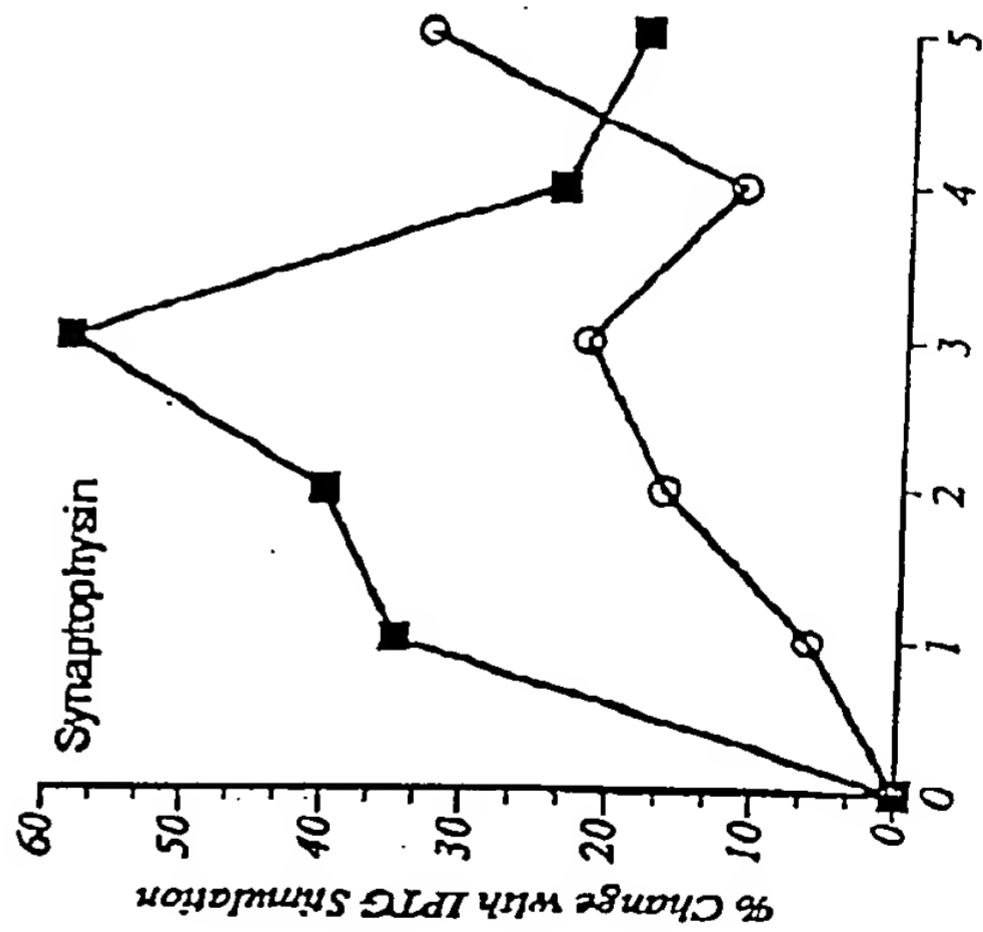
8A



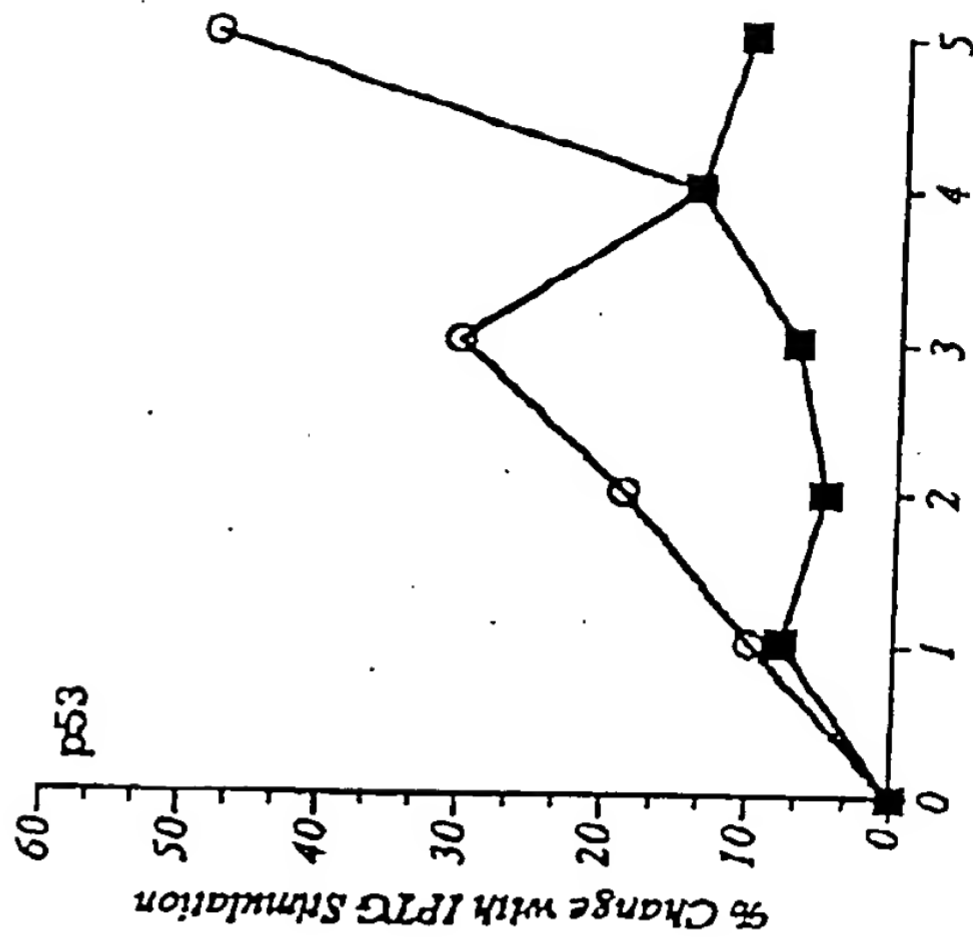
8C

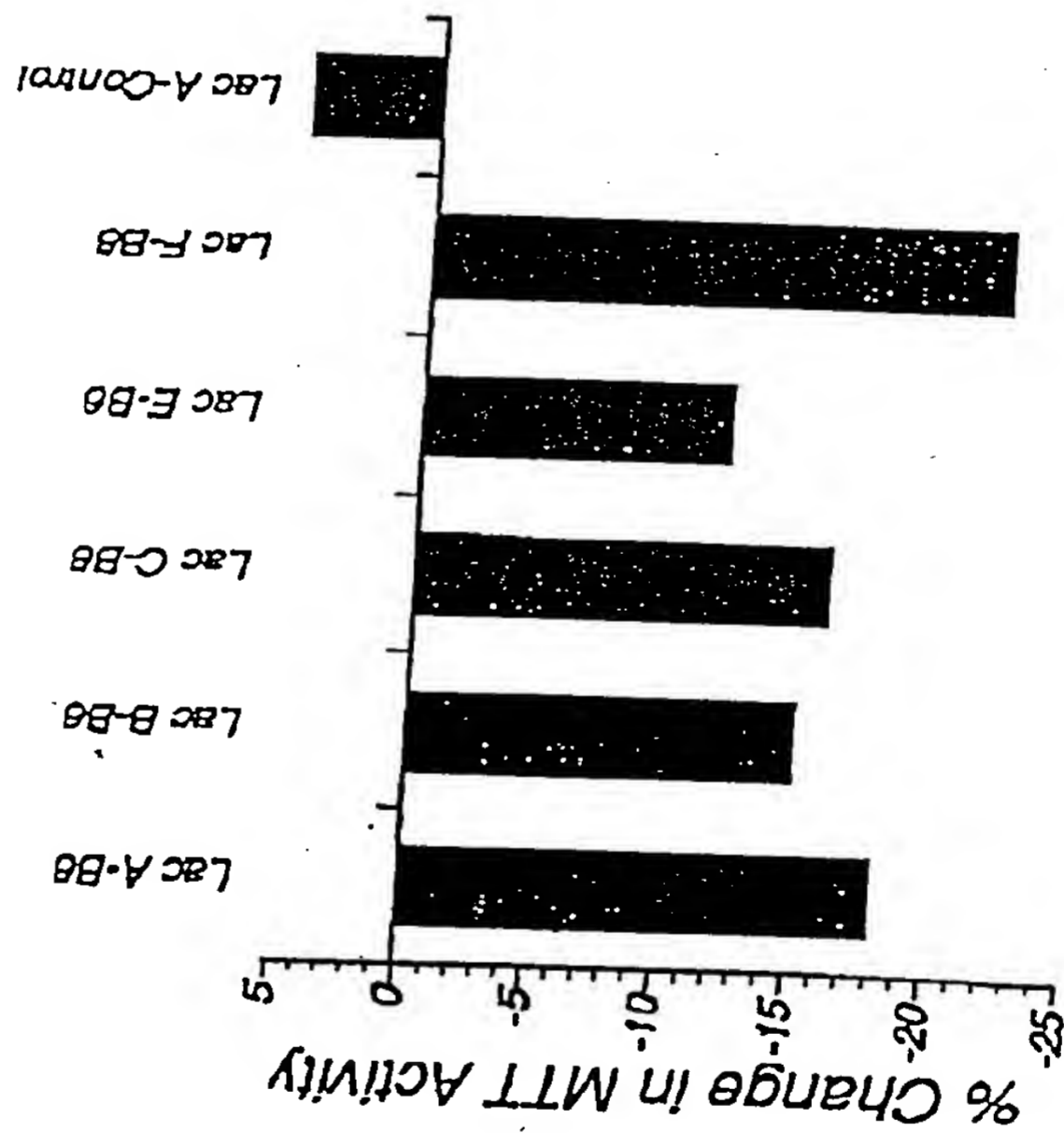
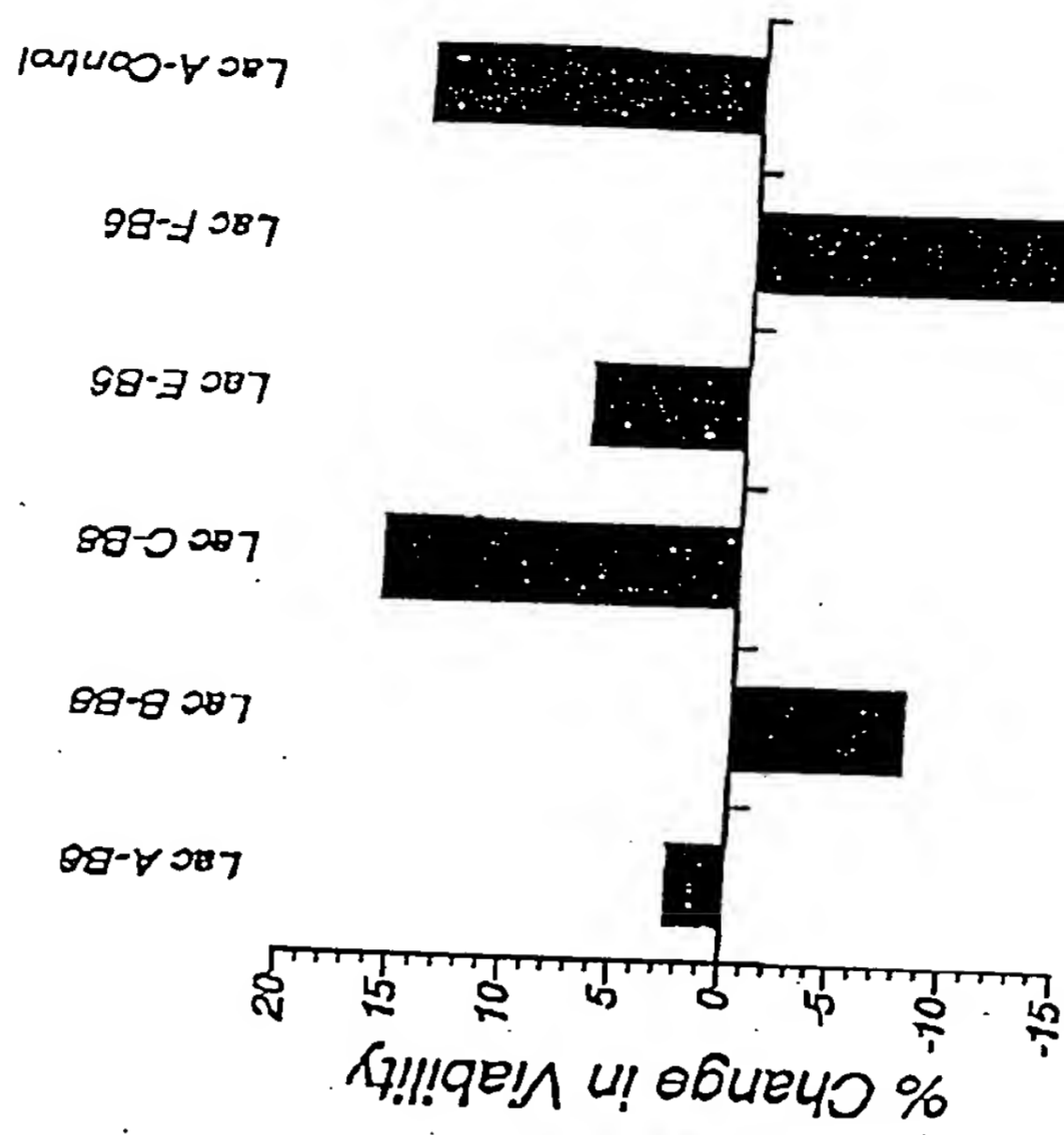


8B



8D

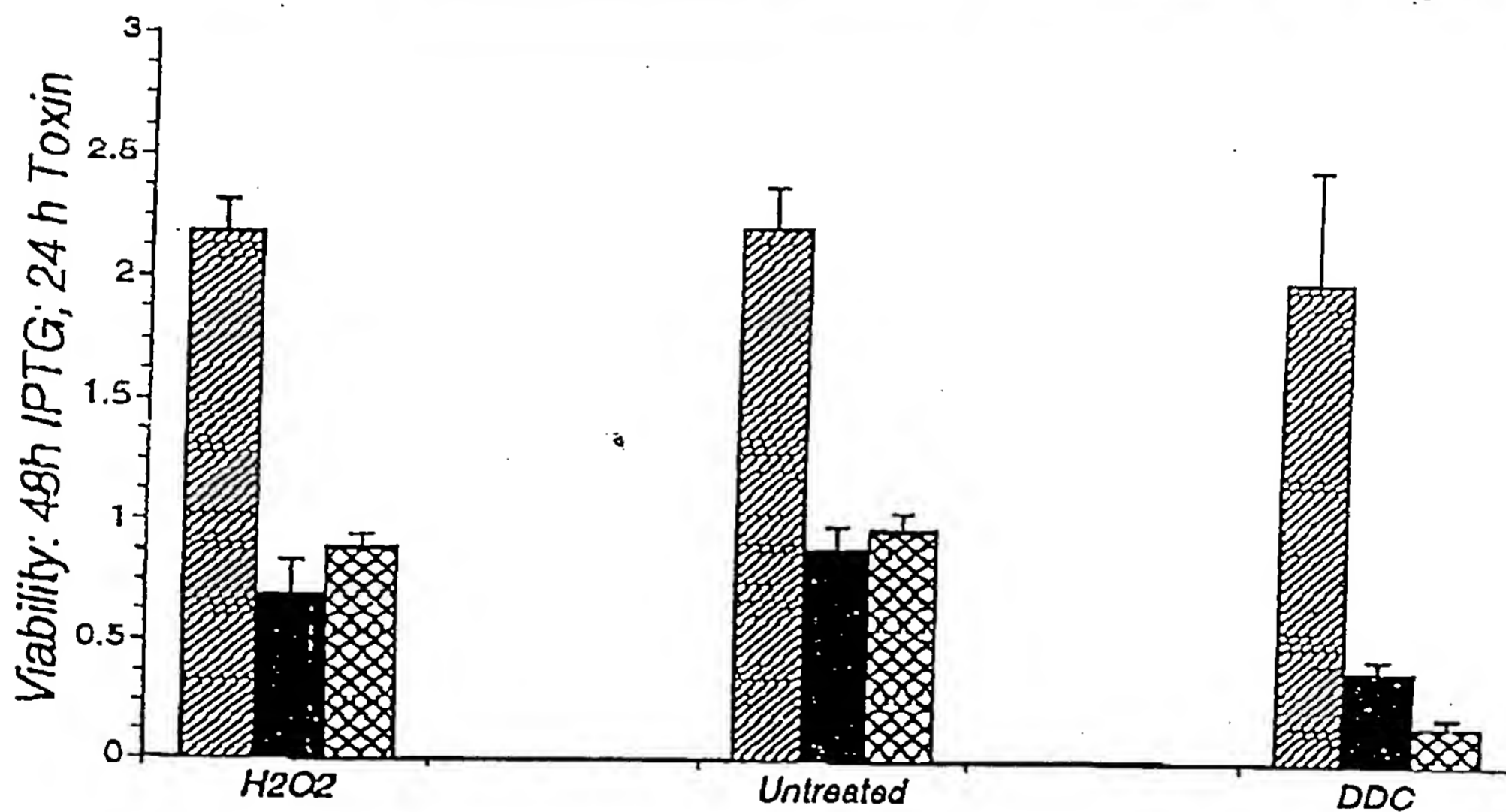
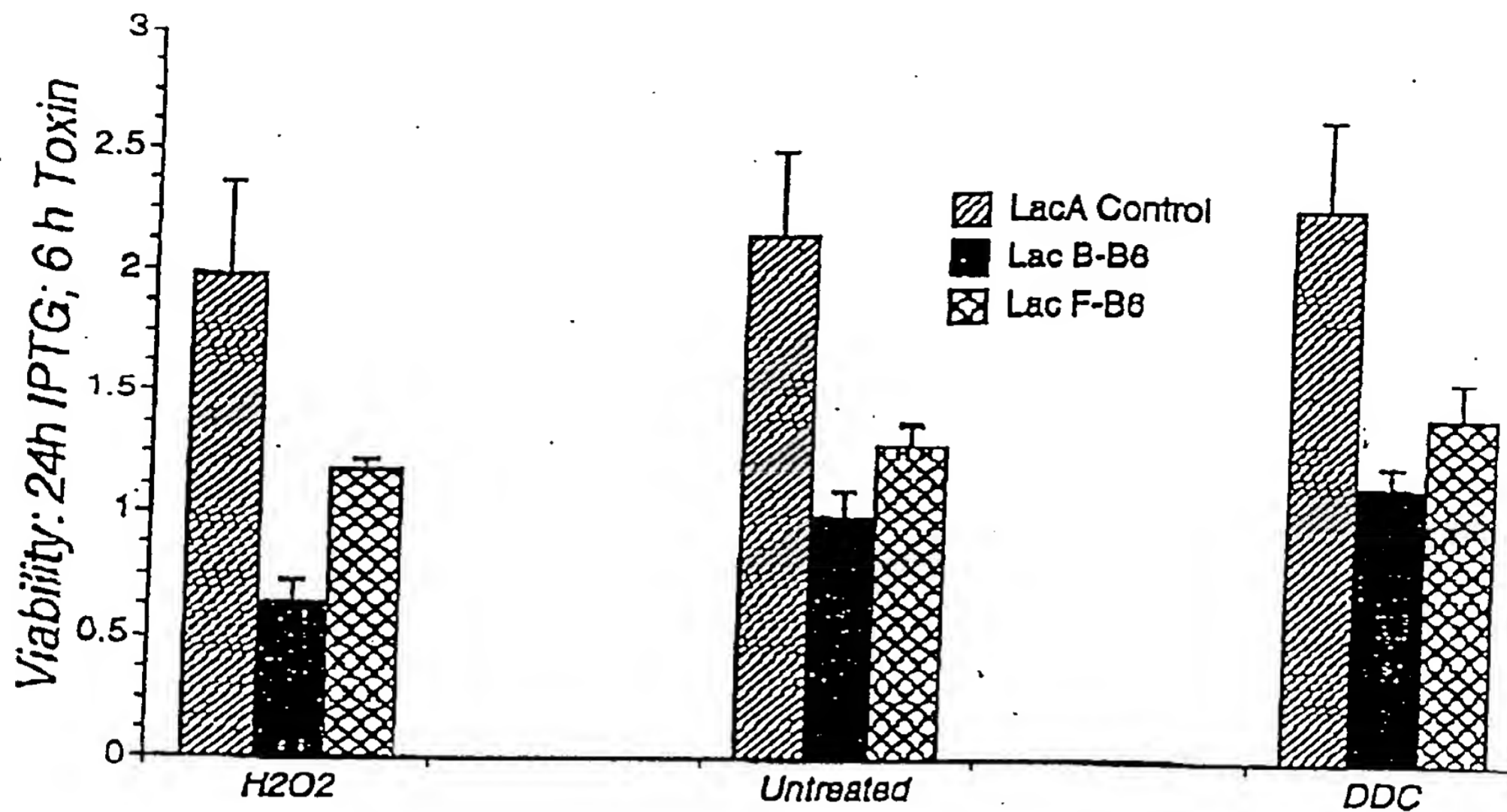




9A

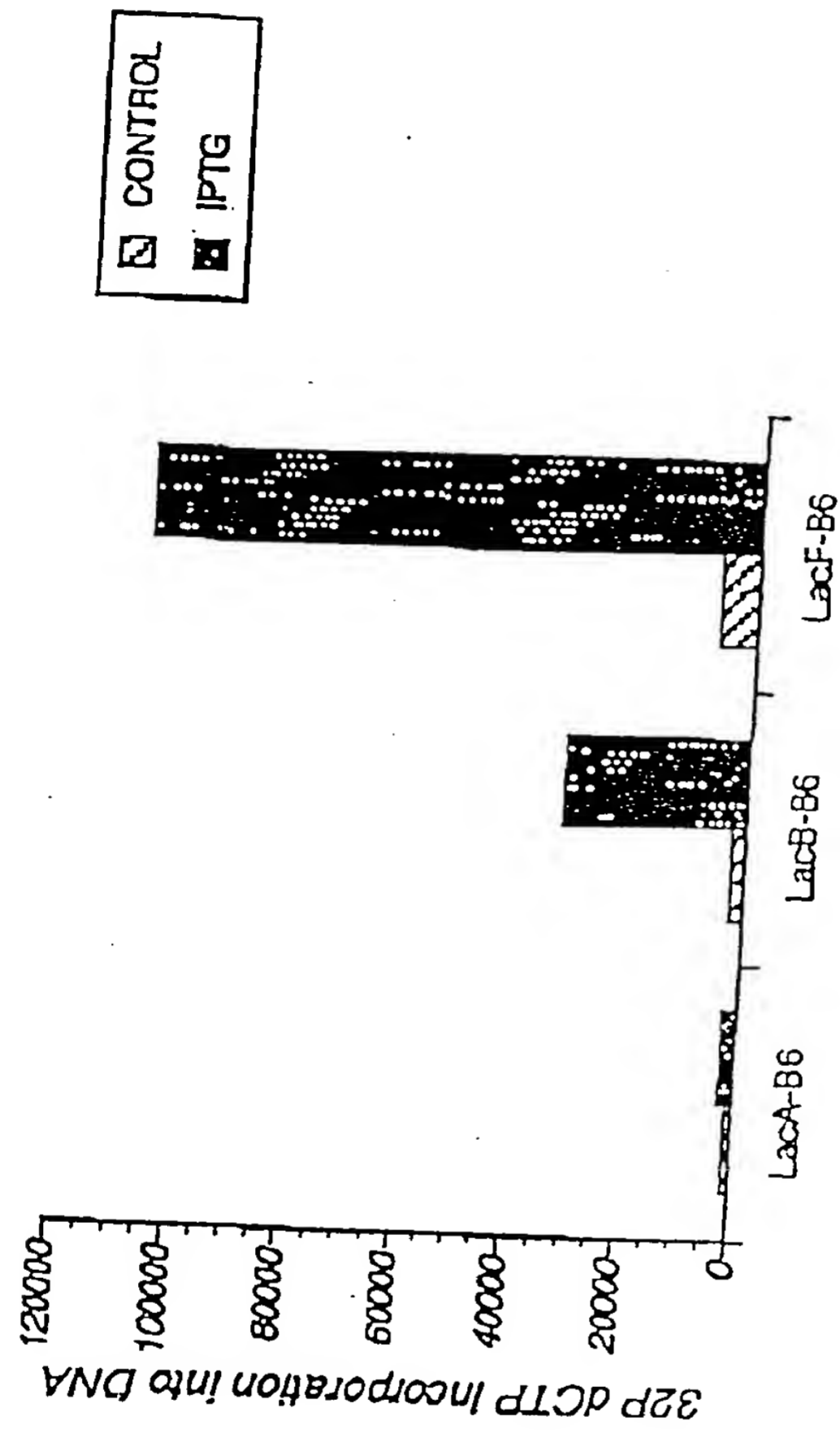
9B

10A



FIGS. 10A-10B

FIG. 11



209260" 2T449660
FIG. 12

